

Applicants' response to the restriction requirement is acknowledged.

However, the sequence listing has not been accepted. It has been determined that that the content of the CRF does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823. As shown on the attached sheet, the issue concerns numeric identifiers <213> and <223>. Apparently, more specific information is required about the source, or identity of the peptide. Applicants should note that this matter is not at the discretion of the examiner, but rather a sequence specialist. In the instant case, the determination of insufficiency was made by Durreshwar Anjum (phone: 866-217-9197). A new CRF listing is required, as well as a paper copy of the sequence listing.



Any inquiry concerning this communication or earlier communications from the examiner should be directed to David Lukton whose telephone number is 571-272-0952. The examiner can normally be reached Monday-Friday from 9:30 to 6:00.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Cecilia Tsang, can be reached at (571)272-0562. The fax number for the organization where this application or proceeding is assigned is 571-273-8300.

Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to the receptionist whose telephone number is 571-272-1600.

/David Lukton/
Primary Examiner, Art Unit 1654

Art Unit: 1654

Attachment to Office action:

Sequence listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9187 (toll free).

Reviewer: Durrasheer Asjua

Timestamp: [year="2009; month="9; day="28; h="11; m="06; sec="03; ms="820;]

Reviewer Comments:

<210> 1

<211> 4

<212> PAT

<213> Artificial Sequence

<220>

<223> peptides

<400> 1

Arg Gly Asp Ser

1

The above is an insufficient response for numeric identifier <213>. Please explain the source of the genetic material. When using "Unknown" for numeric identifier <213>, please provide as much taxonomic information, as possible, about the organism from which the genetic material was extracted. If the genetic material was extracted from a sample in which there was an unknown variety of organisms, please explain where the sample was taken, for example a soil sample. These errors appear in other sequences in the sequence listing. Please make all necessary changes.

Validated By CEFValidator v 1.6.3



Application No: 10593484 Version No: 1.8

Input Set:

Output Set:

Started: 2009-04-05 15:44:57.662
Finished: 2009-04-25 15:56:57.836
Elapsed: 0 hr(s) 0 min(s) 0 sec(ms) 998 ms
Total Warnings: 14
Total Errors: 0
No. of SeqIDs Defined: 19
Actual SeqID Count: 13

Error code	Error Description
* 213	Artificial or Unknown found in <213> in SEQ ID (1)
* 213	Artificial or Unknown found in <213> in SEQ ID (2)
* 213	Artificial or Unknown found in <213> in SEQ ID (3)
* 213	Artificial or Unknown found in <213> in SEQ ID (4)
* 213	Artificial or Unknown found in <213> in SEQ ID (5)
* 213	Artificial or Unknown found in <213> in SEQ ID (6)
* 213	Artificial or Unknown found in <213> in SEQ ID (7)
* 213	Artificial or Unknown found in <213> in SEQ ID (8)
* 213	Artificial or Unknown found in <213> in SEQ ID (9)
* 213	Artificial or Unknown found in <213> in SEQ ID (10)
* 213	Artificial or Unknown found in <213> in SEQ ID (11)
* 213	Artificial or Unknown found in <213> in SEQ ID (12)
* 213	Artificial or Unknown found in <213> in SEQ ID (13)
* 213	Artificial or Unknown found in <213> in SEQ ID (14)